

Additional file 8

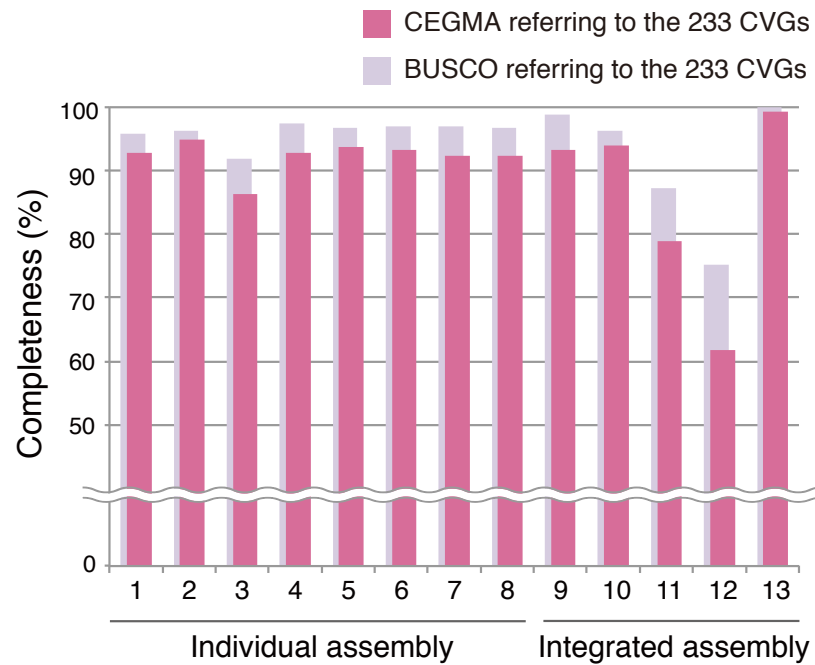


Figure S5. Completeness assessment of transcriptome assemblies by CEGMA and BUSCO referring to CVG.

The scores indicate proportions of the genes recognized as ‘complete’ in individual assemblies by CEGMA and BUSCO in the 233 CVGs. The scores are highly correlated between CEGMA and BUSCO ($R=0.99$; $p=2.3\times 10^{-10}$).